

GenCore version 5.1.6
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OM nucleic - nucleic search, using sw model

Run on: March 16, 2005, 07:53:26 ; Search time 12103 Seconds
(without alignments)
11446.209 Million cell
updates/sec

Title: US-10-616-689-1
Perfect score: 2859
Sequence: 1 atggttatggccctacgac.....tcatctcacaccagccctag
2859

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 4708233 seqs, 24227607955 residues

Total number of hits satisfying chosen parameters: 9416466

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : GenEmbl:
1: gb_ba:
2: gb_htg:
3: gb_in:
4: gb_om:
5: gb_ov:
6: gb_pat:
7: gb_ph:
8: gb_pl:
9: gb_pr:
10: gb_ro:
11: gb_sts:
12: gb_sy:
13: gb_un:
14: gb_vi:

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OM nucleic - nucleic search, using sw model

Run on: March 16, 2005, 07:51:26 ; Search time 1464 Seconds
(without alignments)

11560.478 Million cell
updates/sec

Title: US-10-616-689-1
Perfect score: 2859
Sequence: 1 atggttatggcccctacgac.....tcatctcacaccagccctag
2859

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 4390206 seqs, 2959870667 residues

Total number of hits satisfying chosen parameters: 8780412

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : N_Geneseq_16Dec04:*

1: geneseqn1980s:*

2: geneseqn1990s:*

3: geneseqn2000s:*

4: geneseqn2001as:*

5: geneseqn2001bs:*

6: geneseqn2002as:*

7: geneseqn2002bs:*

8: geneseqn2003as:*

9: geneseqn2003bs:*

10: geneseqn2003cs:*

11: geneseqn2003ds:*

12: geneseqn2004as:*

13: geneseqn2004bs:*

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OM nucleic - nucleic search, using sw model

Run on: March 16, 2005, 08:36:31 ; Search time 8930 Seconds
(without alignments)
12186.534 Million cell
updates/sec

Title: US-10-616-689-1
Perfect score: 2859
Sequence: 1 atggttatggcccctacgac.....tcatctcacaccagccctag
2859

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 34239544 seqs, 19032134700 residues

Total number of hits satisfying chosen parameters: 68479088

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : EST:*

1: gb_est1:*

2: gb_est2:*

3: gb_htc:*

4: gb_est3:*

5: gb_est4:*

6: gb_est5:*

7: gb_est6:*

8: gb_gss1:*

9: gb_gss2:*